

10/663,090

=> d his

(FILE 'HOME' ENTERED AT 16:04:40 ON 28 JAN 2005)

FILE 'REGISTRY' ENTERED AT 16:05:04 ON 28 JAN 2005

L1 SCREEN 1995 AND 2016 AND 2004  
L2 STRUCTURE UPLOADED  
L3 QUE L2 AND L1  
L4 0 S L3  
L5 0 S L3 FULL

FILE 'CAPLUS' ENTERED AT 16:06:05 ON 28 JAN 2005

L6 11311 S (METALLOPHARMAC? OR METAL(P) PHARM? OR RADIOPHARMACEUT?)  
L7 1895 S L6 AND (CHELAT? OR LIGAND?)  
L8 192 S L7 AND (MRI OR MAGNETIC(W) RESONANC? OR NUCLEAR(W) MAGNET? OR NMR)  
L9 117 S L8 AND (CONTRAST(W) AGENT? OR IMAG?)  
L10 41 S L9 AND CONJUGA?  
L11 41 DUP REM L10 (0 DUPLICATES REMOVED)

=> log y

10/663,090

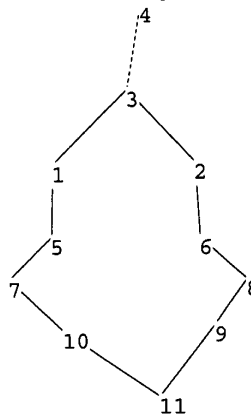
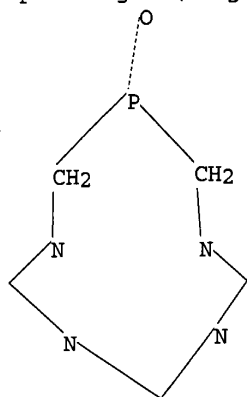
ENTER SCREEN EXPRESSION OR (END):end

=> screen 1995 AND 2016 AND 2004

L1 SCREEN CREATED

=>

Uploading C:\Program Files\Stnexp\Queries\10663090group.str



chain nodes :

4

ring nodes :

1 2 3 5 6 7 8 9 10 11

chain bonds :

3-4

ring bonds :

1-3 1-5 2-3 2-6 5-7 6-8 7-10 8-9 9-11 10-11

exact/norm bonds :

1-3 1-5 2-3 2-6 3-4 5-7 6-8 7-10 8-9 9-11 10-11

Match level :

1:Atom 2:Atom 3:Atom 4:CLASS 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:Atom  
11:Atom

L2 STRUCTURE UPLOADED

=> que L2 AND L1

L3 QUE L2 AND L1

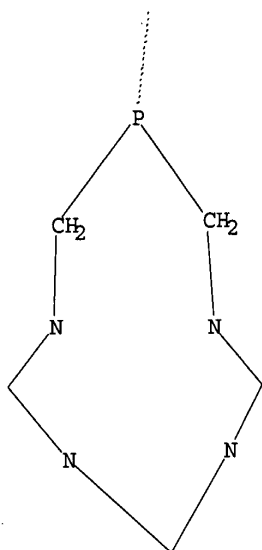
=> d

L3 HAS NO ANSWERS

L1 SCR 1995 AND 2016 AND 2004

L2 STR

10/663,090



Structure attributes must be viewed using STN Express query preparation.  
L3                    QUE ABB=ON PLU=ON L2 AND L1

=> s l3

SAMPLE SEARCH INITIATED 16:05:47 FILE 'REGISTRY'  
SAMPLE SCREEN SEARCH COMPLETED -        2 TO ITERATE

100.0% PROCESSED            2 ITERATIONS                    0 ANSWERS  
SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS:    ONLINE    \*\*COMPLETE\*\*  
                             BATCH    \*\*COMPLETE\*\*  
PROJECTED ITERATIONS:        2 TO        124  
PROJECTED ANSWERS:            0 TO        0

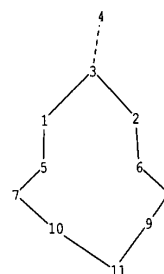
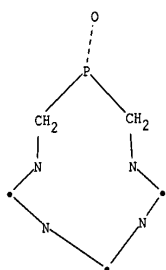
L4                    0 SEA SSS SAM L2 AND L1

=> s l3 full

FULL SEARCH INITIATED 16:05:55 FILE 'REGISTRY'  
FULL SCREEN SEARCH COMPLETED -        27 TO ITERATE

100.0% PROCESSED            27 ITERATIONS                    0 ANSWERS  
SEARCH TIME: 00.00.01

L5                    0 SEA SSS FUL L2 AND L1



chain nodes :

4

ring nodes :

1 2 3 5 6 7 8 9 10 11

chain bonds :

3-4

ring bonds :

1-3 1-5 2-3 2-6 5-7 6-8 7-10 8-9 9-11 10-11

exact/norm bonds :

1-3 1-5 2-3 2-6 3-4 5-7 6-8 7-10 8-9 9-11 10-11

Match level :

1:Atom 2:Atom 3:Atom 4:CLASS 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:Atom 11:Atom